

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/723, 123
Source: IFW0
Date Processed by STIC: 10/26/04

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 10/26/2004

PATENT APPLICATION: US/10/723,123

TIME: 10:48:06

Input Set : A:\Sequence Listing 1998-407-FF-SQ.txt

Output Set: N:\CRF4\10262004\J723123.raw

SEQUENCE LISTING

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4 (1) GENERAL INFORMATION:
C--> 7      (i) APPLICANT: Akzo Nobel N.V.
W--> 15     (ii) TITLE OF INVENTION: Coccidiosis vaccines
17      (iii) NUMBER OF SEQUENCES: 41
W--> 8      (B) STREET: Velperweg 76
9      (C) CITY: Arnhem
10     (E) COUNTRY: The Netherlands
C--> 11     (F) ZIP: 6824 BM
C--> 19     (v) COMPUTER READABLE FORM:
20      (A) MEDIUM TYPE: Floppy disk
21      (B) COMPUTER: IBM PC compatible
22      (C) OPERATING SYSTEM: PC-DOS/MS-DOS
23      (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
C--> 0     (vi) CURRENT APPLICATION DATA:
C--> 0      (A) APPLICATION NUMBER: US/10/723,123
C--> 0      (B) FILING DATE: 26-Nov-2003
C--> 0     (ix) TELECOMMUNICATION INFORMATION:
C--> 12     (A) TELEPHONE: 0412 666379
C--> 13     (B) TELEFAX: 0412 650592
26 (2) INFORMATION FOR SEQ ID NO: 1:
28      (i) SEQUENCE CHARACTERISTICS:
29      (A) LENGTH: 214 amino acids
30      (B) TYPE: amino acid
31      (C) STRANDEDNESS: single
32      (D) TOPOLOGY: linear
34      (ii) MOLECULE TYPE: protein
36      (v) FRAGMENT TYPE: internal
41      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
43 Met  Pro  Phe  Glu  Leu  Pro  Pro  Leu  Pro  Tyr  Pro  Met  Asp  Ala  Leu  Glu
44   1             5             10             15
46 Pro  Tyr  Ile  Ser  Lys  Glu  Thr  Leu  Glu  Tyr  His  Tyr  Gly  Lys  His  His
47   20             25             30
49 Ala  Ala  Tyr  Val  Asn  Asn  Leu  Asn  Arg  Leu  Val  Glu  Gly  Lys  Pro  Glu
50   35             40             45
52 Ala  Ser  Lys  Ser  Leu  Glu  Glu  Ile  Ile  Lys  Thr  Ser  Ser  Gly  Ser  Val
53   50             55             60
55 Leu  Asn  Asn  Ala  Gly  Gln  Ala  Trp  Asn  His  Thr  Phe  Tyr  Trp  Lys  Ser
56  65             70             75             80
58 Met  Arg  Pro  Ala  Ser  Ala  Gly  Gly  Pro  Pro  Gly  Ala  Pro  Gly  Gly  Gly
59   85             90             95
61 Pro  Pro  Gly  Ala  Pro  Gly  Ala  Pro  Leu  Arg  Glu  Glu  Leu  Glu  Ser  Ala
62  100            105            110

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64 Phe Gly Gly Val Glu Lys Phe Arg Glu Ala Phe Ala Ala Ala Ala Ala
65      115                      120                      125
67 Ala His Phe Gly Ser Gly Trp Ala Trp Leu Cys Phe Cys Lys Lys Ser
68      130                      135                      140
70 Arg Ser Leu Phe Leu Leu Gln Thr His Asp Gly Ala Thr Pro Phe Arg
71 145                      150                      155                      160
73 Asp Asn Pro Asn Cys Ala Pro Leu Leu Thr Cys Asp Leu Trp Glu His
74      165                      170                      175
76 Ala Tyr Tyr Ile Asp Arg Arg Asn Asp Arg Lys Ser Tyr Leu Asp Ala
77      180                      185                      190
79 Trp Trp Ser Val Val Asn Trp Asp Phe Ala Asn Glu Asn Leu Lys Lys
80      195                      200                      205
82 Ala Met Gln Gly Ser Asp
83      210
86 (2) INFORMATION FOR SEQ ID NO: 2:
88     (i) SEQUENCE CHARACTERISTICS:
89         (A) LENGTH: 13 amino acids
90         (B) TYPE: amino acid
91         (C) STRANDEDNESS: single
92         (D) TOPOLOGY: linear
94     (ii) MOLECULE TYPE: protein
96     (v) FRAGMENT TYPE: internal
101    (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
103 Leu Gly Pro Leu Ala Leu Pro Leu Leu Ala Asp Val Arg
104      1          5          10
107 (2) INFORMATION FOR SEQ ID NO: 3:
109     (i) SEQUENCE CHARACTERISTICS:
110         (A) LENGTH: 223 amino acids
111         (B) TYPE: amino acid
112         (C) STRANDEDNESS: single
113         (D) TOPOLOGY: linear
115     (ii) MOLECULE TYPE: protein
117     (v) FRAGMENT TYPE: internal
122    (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
124 Met Pro Leu Asn Leu Gly Asp Ser Phe Pro Asp Phe Gln Ala Glu Ala
125      1          5          10          15
127 Leu Gly Ala Glu His Phe Arg Leu His Glu Tyr Leu Gly Asp Ser Trp
128      20          25          30
130 Gly Val Met Phe Ser His Pro Asn Asp Phe Thr Pro Val Cys Thr Thr
131      35          40          45
133 Glu Leu Ala Glu Ala Val Lys Leu Gln Asp Ser Phe Thr Lys Lys Asn
134      50          55          60
136 Cys Lys Leu Val Gly Phe Ser Cys Asn Asp Leu Gln Ser His Arg Glu
137      65          70          75          80
139 Trp Ala Lys Asp Ile Met Ala Tyr Ala Gly Arg Ser Gly Asn Leu Pro
140      85          90          95
142 Phe Pro Leu Val Cys Asp Pro Asn Arg Glu Leu Ala Ala Ser Leu Gly
143      100         105         110
145 Ile Met Asp Pro Ala Glu Lys Asp Lys Lys Gly Leu Pro Leu Thr Cys

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146          115          120          125
148 Arg Cys Val Phe Phe Ile Ser Pro Glu Lys Lys Leu Ala Ala Ser Ile
149      130          135          140
151 Leu Tyr Pro Ala Thr Thr Gly Arg Asn Phe Ala Glu Ile Leu Arg Val
152 145          150          155          160
154 Leu Asp Ser Leu Gln Leu Thr Ala Lys Phe Pro Val Ala Thr Pro Val
155          165          170          175
157 Asp Trp Thr Ala Gly Ala Lys Cys Cys Val Val Pro Asn Leu Ala Ala
158          180          185          190
160 Glu Glu Ala Gln Arg Leu Leu Pro Lys Gly His Glu Ala Leu Gln Leu
161          195          200          205
163 Pro Ser Gly Lys Pro Tyr Leu Arg Leu Thr Pro Asp Pro Arg Gly
164      210          215          220
167 (2) INFORMATION FOR SEQ ID NO: 4:
169     (i) SEQUENCE CHARACTERISTICS:
170         (A) LENGTH: 15 amino acids
171         (B) TYPE: amino acid
172         (C) STRANDEDNESS: single
173         (D) TOPOLOGY: linear
175     (ii) MOLECULE TYPE: protein
177     (v) FRAGMENT TYPE: internal
182     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
184 Met Ser Pro Ser Pro Ala Gly Val Ala Glu Tyr Leu Ala Ser Leu
185 1          5          10          15
187 (2) INFORMATION FOR SEQ ID NO: 5:
189     (i) SEQUENCE CHARACTERISTICS:
190         (A) LENGTH: 16 amino acids
191         (B) TYPE: amino acid
192         (C) STRANDEDNESS: single
193         (D) TOPOLOGY: linear
195     (ii) MOLECULE TYPE: protein
197     (v) FRAGMENT TYPE: internal
202     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
204 Asn His Ala Glu Phe Asp Pro Ser Gln Thr Glu Val Val Val Phe Pro
205 1          5          10          15
208 (2) INFORMATION FOR SEQ ID NO: 6:
210     (i) SEQUENCE CHARACTERISTICS:
211         (A) LENGTH: 20 amino acids
212         (B) TYPE: amino acid
213         (C) STRANDEDNESS: single
214         (D) TOPOLOGY: linear
216     (ii) MOLECULE TYPE: protein
218     (v) FRAGMENT TYPE: internal
223     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
225 Val Asp Ser Phe Thr Pro Ser Val Gly Cys Val Phe Ala Gly Met Pro
226 1          5          10          15
228 Ala Asp Phe Arg
229          20
231 (2) INFORMATION FOR SEQ ID NO: 7:

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233      (i) SEQUENCE CHARACTERISTICS:
234          (A) LENGTH: 17 base pairs
235          (B) TYPE: nucleic acid
236          (C) STRANDEDNESS: double
237          (D) TOPOLOGY: linear
239      (ii) MOLECULE TYPE: DNA (genomic)
244      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
246 GTAAATTGGG ACTTCGC 17
248 (2) INFORMATION FOR SEQ ID NO: 8:
250      (i) SEQUENCE CHARACTERISTICS:
251          (A) LENGTH: 17 base pairs
252          (B) TYPE: nucleic acid
253          (C) STRANDEDNESS: double
254          (D) TOPOLOGY: linear
256      (ii) MOLECULE TYPE: DNA (genomic)
261      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
263 GTAAACTGGG ACTTCGC 17
265 (2) INFORMATION FOR SEQ ID NO: 9:
267      (i) SEQUENCE CHARACTERISTICS:
268          (A) LENGTH: 17 base pairs
269          (B) TYPE: nucleic acid
270          (C) STRANDEDNESS: double
271          (D) TOPOLOGY: linear
273      (ii) MOLECULE TYPE: DNA (genomic)
278      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
280 GTAAATTGGG ACTTCGC 17
282 (2) INFORMATION FOR SEQ ID NO: 10:
284      (i) SEQUENCE CHARACTERISTICS:
285          (A) LENGTH: 17 base pairs
286          (B) TYPE: nucleic acid
287          (C) STRANDEDNESS: double
288          (D) TOPOLOGY: linear
290      (ii) MOLECULE TYPE: DNA (genomic)
295      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
297 GTAAACTGGG ACTTCGC 17
299 (2) INFORMATION FOR SEQ ID NO: 11:
301      (i) SEQUENCE CHARACTERISTICS:
302          (A) LENGTH: 17 base pairs
303          (B) TYPE: nucleic acid
304          (C) STRANDEDNESS: double
305          (D) TOPOLOGY: linear
307      (ii) MOLECULE TYPE: DNA (genomic)
312      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
314 GTTAATTGGG ACTTCGC 17
316 (2) INFORMATION FOR SEQ ID NO: 12:
318      (i) SEQUENCE CHARACTERISTICS:
319          (A) LENGTH: 17 base pairs
320          (B) TYPE: nucleic acid
321          (C) STRANDEDNESS: double

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TIME: 10:48:06

Input Set : A:\Sequence Listing 1998-407-FF-SQ.txt

Output Set: N:\CRF4\10262004\J723123.raw

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322         (D) TOPOLOGY: linear
324     (ii) MOLECULE TYPE: DNA (genomic)
329     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
331 GTTAACTGGG ACTTCGC                                     17
333 (2) INFORMATION FOR SEQ ID NO: 13:
335     (i) SEQUENCE CHARACTERISTICS:
336         (A) LENGTH: 17 base pairs
337         (B) TYPE: nucleic acid
338         (C) STRANDEDNESS: double
339         (D) TOPOLOGY: linear
341     (ii) MOLECULE TYPE: DNA (genomic)
346     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
348 GTTAATTGGG ACTTCGC                                     17
350 (2) INFORMATION FOR SEQ ID NO: 14:
352     (i) SEQUENCE CHARACTERISTICS:
353         (A) LENGTH: 17 base pairs
354         (B) TYPE: nucleic acid
355         (C) STRANDEDNESS: double
356         (D) TOPOLOGY: linear
358     (ii) MOLECULE TYPE: DNA (genomic)
363     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
365 GTTAACTGGG ACTTCGC                                     17
367 (2) INFORMATION FOR SEQ ID NO: 15:
369     (i) SEQUENCE CHARACTERISTICS:
370         (A) LENGTH: 17 base pairs
371         (B) TYPE: nucleic acid
372         (C) STRANDEDNESS: double
373         (D) TOPOLOGY: linear
375     (ii) MOLECULE TYPE: DNA (genomic)
380     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
382 GTGAATTGGG ACTTTGC                                     17
384 (2) INFORMATION FOR SEQ ID NO: 16:
386     (i) SEQUENCE CHARACTERISTICS:
387         (A) LENGTH: 17 base pairs
388         (B) TYPE: nucleic acid
389         (C) STRANDEDNESS: double
390         (D) TOPOLOGY: linear
392     (ii) MOLECULE TYPE: DNA (genomic)
397     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
399 GTGAACTGGG ACTTTGC                                     17
401 (2) INFORMATION FOR SEQ ID NO: 17:
403     (i) SEQUENCE CHARACTERISTICS:
404         (A) LENGTH: 17 base pairs
405         (B) TYPE: nucleic acid
406         (C) STRANDEDNESS: double
407         (D) TOPOLOGY: linear
409     (ii) MOLECULE TYPE: DNA (genomic)
414     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
416 GTGAATTGGG ACTTTGC                                     17

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VERIFICATION SUMMARY

DATE: 10/26/2004

PATENT APPLICATION: US/10/723,123

TIME: 10:48:07

Input Set : A:\Sequence Listing 1998-407-FF-SQ.txt

Output Set: N:\CRF4\10262004\J723123.raw

L:7 M:220 C: Keyword misspelled or invalid format, [(i) APPLICANT:] of (1) line corrected
L:8 M:238 W: Alpha Fields not Ordered, Reordered [(B) STREET:] of (1)(iv)
L:11 M:220 C: Keyword misspelled or invalid format, [(F) ZIP:]
L:12 M:220 C: Keyword misspelled or invalid format, [(A) TELEPHONE:]
L:13 M:220 C: Keyword misspelled or invalid format, [(B) TELEFAX:]
L:15 M:238 W: Alpha Fields not Ordered, Reordered [(ii) TITLE OF INVENTION:] of (1)
L:19 M:220 C: Keyword misspelled or invalid format, [(v) COMPUTER READABLE FORM:]
L:0 M:247 C: Inserted Optional Header Field, [(ix) TELECOMMUNICATION INFORMATION:]
L:0 M:249 C: Inserted Mandatory Field, [(vi) CURRENT APPLICATION DATA:]
L:0 M:249 C: Inserted Mandatory Field, [(A) APPLICATION NUMBER:]
L:0 M:249 C: Inserted Mandatory Field, [(B) FILING DATE:]
L:790 M:111 C: (47) String data converted to upper case,
L:794 M:111 C: (47) String data converted to upper case,
L:798 M:111 C: (47) String data converted to upper case,
L:802 M:111 C: (47) String data converted to upper case,
L:806 M:111 C: (47) String data converted to upper case,
L:810 M:111 C: (47) String data converted to upper case,
L:814 M:111 C: (47) String data converted to upper case,
L:818 M:111 C: (47) String data converted to upper case,
L:822 M:111 C: (47) String data converted to upper case,
L:826 M:111 C: (47) String data converted to upper case,
L:830 M:111 C: (47) String data converted to upper case,
L:834 M:111 C: (47) String data converted to upper case,
L:838 M:111 C: (47) String data converted to upper case,
L:842 M:111 C: (47) String data converted to upper case,
L:844 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:39
L:846 M:111 C: (47) String data converted to upper case,
L:890 M:111 C: (47) String data converted to upper case,
L:894 M:111 C: (47) String data converted to upper case,
L:898 M:111 C: (47) String data converted to upper case,
L:902 M:111 C: (47) String data converted to upper case,
L:906 M:111 C: (47) String data converted to upper case,
L:910 M:111 C: (47) String data converted to upper case,
L:914 M:111 C: (47) String data converted to upper case,
L:918 M:111 C: (47) String data converted to upper case,
L:922 M:111 C: (47) String data converted to upper case,
L:926 M:111 C: (47) String data converted to upper case,
L:930 M:111 C: (47) String data converted to upper case,
L:934 M:111 C: (47) String data converted to upper case,
L:938 M:111 C: (47) String data converted to upper case,
L:942 M:111 C: (47) String data converted to upper case,